

*A 1 cont'd*

4:1288-1303, 1990) with the *Hind*III-containing primers *aagctt*CTGGCCTTCGAGAACTCCTGCACC (upper strand, *Hind*III in lower case, SEQ ID NO: 25) and *aagctt*AGTGCCACGTGAAGGAGCGAGGCC (lower strand, SEQ ID NO: 26). The resulting 272 bp fragment, which corresponds to the putative catalytic domain of FUT3 (*Id.*), was cloned in reverse orientation into the *Hind*III site of pcDNA3 to yield the vector pcDNA3-FUT3CD AS (data not shown).

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